



00670658.TXT

## SEQUENCE LISTING

<110> Chu, Rea-Min  
Lin, Ching-Yi  
Hsiao, Ya-Wen  
Liao, Kuang-Wen

<120> COMPLEX IMMUNO-GENE MEDICAL COMPOSITION FOR INHIBITING TUMOR CELLS

<130> P/741-176

<140> 10/798,096

<141> 2004-03-11

<160> 4

<170> PatentIn version 3.2

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Gly Leu Leu Leu Val Leu Pro Ala Ala Phe Pro Ala Pro Val Pro Pro	
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Gly Glu Asp Ser Lys Asp Val Ala Ala Pro His Arg Gln Pro Leu Thr	
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tct tca gaa cga att gac aaa caa att cgg tac atc ctc gac ggc atc	192
Ser Ser Glu Arg Ile Asp Lys Gln Ile Arg Tyr Ile Leu Asp Gly Ile	
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Ser Ala Leu Arg Lys Glu Thr Cys Asn Lys Ser Asn Met Cys Glu Ser	
65 70 75 80	
agc aaa gag gca ctg gca gaa aac aac ctg aac ctt cca aag atg gct	288
Ser Lys Glu Ala Leu Ala Glu Asn Asn Leu Asn Leu Pro Lys Met Ala	
85 90 95	
gaa aaa gat gga tgc ttc caa tct gga ttc aat gag gag act tgc ctg	336
Glu Lys Asp Gly Cys Phe Gln Ser Gly Phe Asn Glu Glu Thr Cys Leu	
100 105 110	
gtg aaa atc atc act ggt ctt ttg gag ttt gag gta tac cta cag tac	384
Val Lys Ile Ile Thr Gly Leu Leu Glu Phe Glu Val Tyr Leu Gln Tyr	
115 120 125	
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Met	Ser	Thr	Lys	Val	Leu	Ile	Gln	Phe	Leu	Gln	Lys	Lys	Ala	Lys	Asn	160	
145					150					155							
cta	gat	gca	ata	acc	cct	gac	cca	acc	aca	aat	gcc	agc	ctg	ctg		528	
Leu	Asp	Ala	Ile	Thr	Pro	Asp	Pro	Thr	Thr	Asn	Ala	Ser	Leu	Leu		175	
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Thr	Lys	Leu	Gln	Ala	Gln	Asn	Gln	Trp	Leu	Gln	Asp	Met	Thr	Thr	His		
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Leu	Ile	Leu	Arg	Ser	Phe	Lys	Glu	Phe	Leu	Gln	Ser	Ser	Leu	Arg	Ala		
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1				5				10						15			
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Gln	Ser	Met	His	Ile	Asp	Ala	Thr	Leu	Tyr	Thr	Glu	Ser	Asp	Val	His		
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ccc	agt	tgc	aaa	gta	aca	gca	atg	aag	tgc	ttt	ctc	ttg	gag	tta	caa	144	
Pro	Ser	Cys	Lys	Val	Thr	Ala	Met	Lys	Cys	Phe	Leu	Leu	Glu	Leu	Gln		

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aca gaa tct gga tgc aaa gaa tgt gag gaa ctg gag gaa aaa aat att Thr Glu Ser Gly Cys Lys Glu Cys Glu Glu Leu Glu Glu Lys Asn Ile 85 90 95			288
aaa gaa ttt ttg cag agt ttt gta cat att gtc caa atg ttc atc aac Lys Glu Phe Leu Gln Ser Phe Val His Ile Val Gln Met Phe Ile Asn 100 105 110			336
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gtc aca aac agt aac tgg gtg aat gta ata agt gat ttg aaa aaa att Val Thr Asn Ser Asn Trp Val Asn Val Ile Ser Asp Leu Lys Lys Ile 20 25 30			96
gaa gat ctt att caa tct atg cat att gat gct act tta tat acg gaa Glu Asp Leu Ile Gln Ser Met His Ile Asp Ala Thr Leu Tyr Thr Glu 35 40 45			144
agt gat gtt cac ccc agt tgc aaa gta aca gca atg aag tgc ttt ctc Ser Asp Val His Pro Ser Cys Lys Val Thr Ala Met Lys Cys Phe Leu 50 55 60			192
ttg gag tta caa gtt att tca ctt gag tcc gga gat gca agt att cat Leu Glu Leu Gln Val Ile Ser Leu Glu Ser Gly Asp Ala Ser Ile His 65 70 75 80			240
gat aca gta gaa aat ctg atc atc cta gca aac aac agt ttg tct tct Asp Thr Val Glu Asn Leu Ile Ile Leu Ala Asn Asn Ser Leu Ser Ser 85 90 95			288
aat ggg aat gta aca gaa tct gga tgc aaa gaa tgt gag gaa ctg gag Asn Gly Asn Val Thr Glu Ser Gly Cys Lys Glu Cys Glu Glu Leu Glu 100 105 110			336
gaa aaa aat att aaa gaa ttt ttg cag agt ttt gta cat att gtc caa Glu Lys Asn Ile Lys Glu Phe Leu Gln Ser Phe Val His Ile Val Gln 115 120 125			384

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atg ttc atc aac act tct  
Met Phe Ile Asn Thr Ser  
130

402